

IN THE CLAIMS:

Please cancel Claim 3 without prejudice to or disclaimer of the subject matter presented therein. Please amend Claims 2 and 4 as shown below.

1. (Cancelled)

2. (Currently Amended) A probe set comprising multiple probes that are isolated and that can be used for identification of an HLA-A allele contained in a specimen ~~under a condition in which at least one of the multiple probes specifically hybridizes to the HLA-A allele, wherein each of the multiple probes is a partial sequence of 10 to 30 successive bases of a sequence of an allele in the allele list for HLA-A in the specification, the partial sequence containing a base represented by a capital letter, and wherein the multiple probes, as taken all together and represented in small and capital letters as in the allele list for HLA-A, contain all the bases represented by capital letters in the allele list for HLA-A wherein the multiple probes comprise the probes of SEQ ID NOs. 251 to 454 or the probes of SEQ ID NOs. 455 to 631 so that the 250 alleles of A*010101 through A*8001 listed in the allele list for HLA-A~~ SEQ ID NOs. 1 to 250 can be identified by conducting a set of polymerase chain reaction (PCR) operations using different probes selected from the probe set based on the allele-probe correspondence list shown in Tables 3-1 to 3-9 or in Tables 4-1 to 4-9.

3. (Cancelled)

4. (Withdrawn-Currently Amended) A method for identification of an HLA-A allele contained in a sample from a specimen ~~using a probe set, wherein the probe set is~~ comprising conducting a set of PCR operations using different probes selected from the probe set according to claim 2 ~~or 3~~.

5 to 22. (Cancelled)